Johnson

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RAW SEQUENCE LISTING PATENT APPLICATION US/09/077,214

DATE: 03/03/1999 TIME: 14:38:42

INPUT SET: S30901.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

	1				SEQUENCE	LISTING	
	2						
	3	(1) G	eneral Info	rmation:			ENTERED
	4	(2)	1001 TO11	0-1	v-14		
	5	(1)	APPLICANT:	Birnstiel			
	6 7				ffer, Tamas		
	8			Steinlein			
	9			Buschle,			
	10			Dubellie, !	ichaci		
	11	(ii)	TITLE OF I	NVENTION: '	rumor Vaccin	e And Process Fo	or the
	12	(,		tion There			
	13		-				
	14	(iii)	NUMBER OF	SEQUENCES:	33		
	15						
	16	(iv)	CORRESPOND				
	17		(A) ADDRE	SSEE: Ster	ne, Kessler,	Goldstein & For	R.L.L.C.
	18				w York Avenu	e N.W.	
	19			Washington	1		
	20		(D) STATE				
	21		, ,	RY: U.S.A.			
	22		(F) ZIP:	20005			
	23	()	COMPLIMED D		D14 -		
	24 25	(V)	COMPUTER RI	M TYPE: Flo			
	25 26				oppy disk C compatible		
	26 27				M: PC-DOS/MS		
	28					#1.0, Version #1	1.30
	29		(D) 501 111	nin. Iuccii	JIII NOICUDO	112.07 (0252012 112	
	30	(ví)	CURRENT AP	PLICATION I	DATA:		
~->	31	,,		CATION NUM		assigned	
	32		(B) FILING	G DATE:		•	
	33		(C) CLASS	IFICATION:			
	34						
	35	(vii)	PRIOR APPL	ICATION DAT	ΓA:		
	36		, ,		BER: DE 195	43 649.0	
	37		(B) FILING	G DATE: 23	-NOV-1995		
	38						
	39	(Vli)	PRIOR APPL				
	40		• •		BER: DE 196	07 044.9	
	41 42		(B) LITITIO	G DATE: 24-	-LEB-1330		
	43	(3ri i i \	ATTORNEY/AG	בפאיי דאופירסא	ANTION.		
	44	(^ T T T)	•	Fleshner,			
	45				MBER: 34,331		
	46				NUMBER: 06	52.1710000	

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	(ix) TELECOMMUNICATION INFORMATION:
	(A) TELEPHONE: 202-371-2600
	(B) TELEFAX: 202-371-2540
(2)	INFORMATION FOR SEQ ID NO:1:
	(i) SEQUENCE CHARACTERISTICS:
	(A) LENGTH: 9 amino acids
	(B) TYPE: amino acid
	(C) STRANDEDNESS: single
	(D) TOPOLOGY: Not Relevant
	(ii) MOLECULE TYPE: peptide
	• •
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
	(-12, 22, 23, 23, 23, 23, 23, 23, 23, 23, 2
	Leu Phe Glu Ala Ile Glu Gly Phe Ile
	1 5
	_
(2)	INFORMATION FOR SEO ID NO:2:
,-,	~ · · · · · · · · · · · · · · · · · · ·
	(i) SEQUENCE CHARACTERISTICS:
	(A) LENGTH: 9 amino acids
	(B) TYPE: amino acid
	(C) STRANDEDNESS: single
	(D) TOPOLOGY: Not Relevant
	(0, 1010_001, 1.00 1.010 1.010
	(ii) MOLECULE TYPE: peptide
	(15) 110 110 111 popular
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
	(112, 412, 112, 112, 112, 112, 112, 112,
	Phe Phe Ile Gly Ala Leu Glu Glu Ile
	1 5
(2)	INFORMATION FOR SEQ ID NO:3:
(-,	
	(i) SEQUENCE CHARACTERISTICS:
	(A) LENGTH: 9 amino acids
	(B) TYPE: amino acid
	(C) STRANDEDNESS: single
	(D) TOPOLOGY: Not Relevant
	,_,
	(ii) MOLECULE TYPE: peptide
	.
	(2)

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100	
101	
102	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
103	
104	Ala Ser Asn Glu Asn Met Glu Thr Met
105	1 5
106	
107	(2) INFORMATION FOR SEQ ID NO:4:
108	(2) 2112 0111111 2111 212 == -1111
109	(i) SEQUENCE CHARACTERISTICS:
110	(A) LENGTH: 9 amino acids
111	(B) TYPE: amino acid
112	(C) STRANDEDNESS: single
	(D) TOPOLOGY: Not Relevant
113	(D) TOPOLOGI: NOC RETEVALLE
114	(''') WOTTOTTE MUDE, markido
115	(ii) MOLECULE TYPE: peptide
116	
117	
118	
119	
120	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
121	
122	Leu Pro Glu Ala Ile Glu Gly Phe Gly
123	1 5
124	
125	(2) INFORMATION FOR SEQ ID NO:5:
126	(2)
127	(i) SEQUENCE CHARACTERISTICS:
128	(A) LENGTH: 9 amino acids
129	(B) TYPE: amino acid
130	(C) STRANDEDNESS: single
131	(D) TOPOLOGY: Not Relevant
132	(b) Totoboot. Not notovano
133	(ii) MOLECULE TYPE: peptide
	(II) MODECODE TIPE. Peptide
134	
135	
136	
137	/ !) charged become on the Mo. C.
138	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
139	
140	Ser Pro Ser Tyr Val Tyr His Gln Phe
141	1 5
142	
143	(2) INFORMATION FOR SEQ ID NO:6:
144	
145	(i) SEQUENCE CHARACTERISTICS:
146	(A) LENGTH: 8 amino acids
147	(B) TYPE: amino acid
148	(C) STRANDEDNESS: single
149	(D) TOPOLOGY: Not Relevant
150	
151	(ii) MOLECULE TYPE: peptide
131	(II) MODECODE IIFE. Pepciae
152	(11) MODECODE 11FE. peptide

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153		
154		
155		
156		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
		(XI) DEGOLIGE DEDOCATITIONS DEG TO 10.0.
157		m1 'm1 m1 m
158		Phe Glu Gln Asn Thr Ala Gln Ala
159		1 5
160		
161	(2)	INFORMATION FOR SEQ ID NO:7:
162	,	•
163		(i) SEQUENCE CHARACTERISTICS:
		(A) LENGTH: 8 amino acids
164		
165		(B) TYPE: amino acid
166		(C) STRANDEDNESS: single
167		(D) TOPOLOGY: Not Relevant
168		
169		(ii) MOLECULE TYPE: peptide
170		(12) Manager Page 1
171		
172		
173		
174		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
175		
176		Phe Glu Gln Asn Thr Ala Gln Pro
177		1 5
		1 3
178	(0)	TYPODWARTON FOR GEO TO NO. C.
179	(2)	INFORMATION FOR SEQ ID NO:8:
180		
181		(i) SEQUENCE CHARACTERISTICS:
182		(A) LENGTH: 9 amino acids
183		(B) TYPE: amino acid
184		(C) STRANDEDNESS: single
185		(D) TOPOLOGY: Not Relevant
		(5) 10102001. 1100 1102014110
186		(ii) war naur mune, mantida
187		(ii) MOLECULE TYPE: peptide
188		
189		
190		
191		
192		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
193		,,,
194		Ser Tyr Phe Pro Glu Ile Thr His Ile
195		1 5
196		
197	(2)	INFORMATION FOR SEQ ID NO:9:
198		
199		(i) SEQUENCE CHARACTERISTICS:
200		(A) LENGTH: 9 amino acids
201		(B) TYPE: amino acid
202		(C) STRANDEDNESS: single
202		(D) TOPOLOGY: Not Relevant
		(D) TOPOHOGI. NOC RETEVANC
204		(11) area novem novem novembrida
205		(ii) MOLECULE TYPE: peptide

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206
207
208
209
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
210
211
          Glu Ala Asp Pro Thr Gly His Ser Tyr
212
213
214
     (2) INFORMATION FOR SEQ ID NO:10:
215
216
           (i) SEQUENCE CHARACTERISTICS:
217
                (A) LENGTH: 9 amino acids
218
                (B) TYPE: amino acid
219
                (C) STRANDEDNESS: single
220
                (D) TOPOLOGY: Not Relevant
221
222
223
         (ii) MOLECULE TYPE: peptide
224
225
226
227
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
228
229
          Glu Val Asp Pro Ile Gly His Leu Tyr
230
                           5
231
          1
232
     (2) INFORMATION FOR SEQ ID NO:11:
233
234
           (i) SEQUENCE CHARACTERISTICS:
235
                (A) LENGTH: 9 amino acids
236
237
                (B) TYPE: amino acid
                (C) STRANDEDNESS: single
238
                (D) TOPOLOGY: Not Relevant
239
240
          (ii) MOLECULE TYPE: peptide
241
242
243
244
245
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:
246
247
           Tyr Met Asn Gly Thr Met Ser Gln Val
248
                           5
249
           1
250
      (2) INFORMATION FOR SEQ ID NO:12:
251
252
           (i) SEQUENCE CHARACTERISTICS:
253
                (A) LENGTH: 9 amino acids
254
255
                (B) TYPE: amino acid
                (C) STRANDEDNESS: single
256
                (D) TOPOLOGY: Not Relevant
257
258
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SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/077,214

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Line

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Error

Wrong application Serial Number

Original Text

(A) APPLICATION NUMBER: To be assigned